RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:

Source:

Date Processed by STIC:

ENTERED



TFWO

RAW SEQUENCE LISTING DATE: 02/20/2007 PATENT APPLICATION: US/10/576,906 TIME: 14:53:50

Input Set : A:\19781.seq.txt

Output Set: N:\CRF4\02152007\J576906.raw

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3 <110> APPLICANT: BARTHOLOMEUSZ, Angeline; Ingrid
        LOCARNINI, Stephen, Alister
 4
        AYRES, Anna
 5
        YUEN, Lilly, Ka, Wai
        SASADEUSZ, Joseph, John
 9 <120> TITLE OF INVENTION: HBV VARIANTS DETECTION AND APPLICATION
                                                                       P5.6)
11 <130> FILE REFERENCE: 19781
13 <140> CURRENT APPLICATION NUMBER: 10/576,906
14 <141> CURRENT FILING DATE: 2006-04-21
16 <150> PRIOR APPLICATION NUMBER: AU2003905776
17 <151> PRIOR FILING DATE: 2003-10-21
19 <150> PRIOR APPLICATION NUMBER: AU2004900962
20 <151> PRIOR FILING DATE: 2004-02-25
22 <160> NUMBER OF SEQ ID NOS: 42
24 <170> SOFTWARE: PatentIn version 3.2
26 <210> SEQ ID NO: 1
27 <211> LENGTH: 1007
28 <212> TYPE: DNA
29 <213> ORGANISM: hepatitis B virus
31 <400> SEQUENCE: 1
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36 agacctgctg gtggctccag ttccggaaca gtaaaccctg ttccgactac tgcctctccc
38 atategteaa tettetegag gaetggggae eetgegeega atatggagag caccacatea
                                                                         240
                                                                         300
40 ggattcctag gacccctgct cgtgttacag gcggggtttt tcttgttgac aagaatcctc
                                                                        360
42 acaataccaa agagtctaga ctcgtggtgg acttctctca attttctagg gggagcaccc
44 acqtqtcctq qccaaaattt qcaqtcccca acctccaatc actcaccaac ctcttqtcct
                                                                        420
46 ccaatttgtc ctggttatcg ctggatgtgt ctgcggcgtt ttatcatctt cctcttcatc
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48 ctgctgctat gcctcatctt cttgttggtt cttctggact accaaggtat gttgcccgtt
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50 tgtcctctac ttccaggaac atcaactacc agcacgggac catgcaagac ctgcacgact
                                                                        600
52 cctgctcaag gaacctctat gtttccctct tgttgctgta caaaaccttc ggacggaaat
                                                                        660
54 tgcacttgta ttcccatccc atcatcttgg gctttcgtaa gattcctatg ggagtgggcc
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56 tragtrogtt totootggtt cagtttacta gtgccatttg ttcagtggtt cgtagggctt
                                                                        780
58 tececeactg tttggettte agttatatgg atgatgtggt attgggggee aagtetgtae
                                                                         840
60 aacatcttga atccctttat accgctatta ccaattttct tttgtctttg ggtatacatt
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62 taaaccctaa taaaaccaag cgttggggct actcccttaa cttcatggga tatgtaattg
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68 <211> LENGTH: 270
69 <212> TYPE: PRT
70 <213> ORGANISM: hepatitis B virus
72 <400> SEQUENCE: 2
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74 Glu Asp Trp Gly Pro Cys Ala Glu Tyr Gly Glu His His Ile Arg Ile

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75 1 10 78 Pro Arg Thr Pro Ala Arg Val Thr Gly Gly Val Phe Leu Val Asp Lys 20 82 Asn Pro His Asn Thr Lys Glu Ser Arg Leu Val Val Asp Phe Ser Gln 86 Phe Ser Arg Gly Ser Thr His Val Ser Trp Pro Lys Phe Ala Val Pro 55 90 Asn Leu Gln Ser Leu Thr Asn Leu Leu Ser Ser Asn Leu Ser Trp Leu 70 94 Ser Leu Asp Val Ser Ala Ala Phe Tyr His Leu Pro Leu His Pro Ala 90 98 Ala Met Pro His Leu Leu Val Gly Ser Ser Gly Leu Pro Arg Tyr Val 105 100 110 102 Ala Arg Leu Ser Ser Thr Ser Arg Asn Ile Asn Tyr Gln His Gly Thr 120 106 Met Gln Asp Leu His Asp Ser Cys Ser Arg Asn Leu Tyr Val Ser Leu 135 110 Leu Leu Tyr Lys Thr Phe Gly Arg Lys Leu His Leu Tyr Ser His 114 Pro Ile Ile Leu Gly Phe Arg Lys Ile Pro Met Gly Val Gly Leu Ser 170 165 118 Pro Phe Leu Leu Val Gln Phe Thr Ser Ala Ile Cys Ser Val Val Arg 185 122 Arg Ala Phe Pro His Cys Leu Ala Phe Ser Tyr Met Asp Asp Val Val 195 200 126 Leu Gly Ala Lys Ser Val Gln His Leu Glu Ser Leu Tyr Thr Ala Ile 210 215 130 Thr Asn Phe Leu Leu Ser Leu Gly Ile His Leu Asn Pro Asn Lys Thr 230 235 134 Lys Arg Trp Gly Tyr Ser Leu Asn Phe Met Gly Tyr Val Ile Gly Ser 250 245 138 Trp Gly Thr Leu Pro Gln Glu His Ile Val Gln Lys Ile Lys 260 265 142 <210> SEQ ID NO: 3 143 <211> LENGTH: 226 144 <212> TYPE: PRT 145 <213> ORGANISM: hepatitis B virus 147 <400> SEQUENCE: 3 149 Met Glu Ser Thr Thr Ser Gly Phe Leu Gly Pro Leu Leu Val Leu Gln 10 153 Ala Gly Phe Phe Leu Leu Thr Arg Ile Leu Thr Ile Pro Lys Ser Leu 20 25 157 Asp Ser Trp Trp Thr Ser Leu Asn Phe Leu Gly Gly Ala Pro Thr Cys 40 161 Pro Gly Gln Asn Leu Gln Ser Pro Thr Ser Asn His Ser Pro Thr Ser 165 Cys Pro Pro Ile Cys Pro Gly Tyr Arg Trp Met Cys Leu Arg Arg Phe 169 Ile Ile Phe Leu Phe Ile Leu Leu Cys Leu Ile Phe Leu Leu Val RAW SEQUENCE LISTING DATE: 02/20/2007
PATENT APPLICATION: US/10/576,906 TIME: 14:53:50

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170
173 Leu Leu Asp Tyr Gln Gly Met Leu Pro Val Cys Pro Leu Leu Pro Gly
               100
                                  105
177 Thr Ser Thr Thr Ser Thr Gly Pro Cys Lys Thr Cys Thr Thr Pro Ala
                              120
                                                 125
178
           115
181 Gln Gly Thr Ser Met Phe Pro Ser Cys Cys Cys Thr Lys Pro Ser Asp
                          135
                                             140
185 Gly Asn Cys Thr Cys Ile Pro Ile Pro Ser Ser Trp Ala Phe Val Arg
                      150
                                          155
189 Phe Leu Trp Glu Trp Ala Ser Val Arg Phe Ser Trp Phe Ser Leu Leu
                  165
                                      170
193 Val Pro Phe Val Gln Trp Phe Val Gly Leu Ser Pro Thr Val Trp Leu
                                  185
197 Ser Val Ile Trp Met Met Trp Tyr Trp Gly Pro Ser Leu Tyr Asn Ile
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201 Leu Asn Pro Phe Ile Pro Leu Leu Pro Ile Phe Phe Cys Leu Trp Val
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205 Tyr Ile
206 225
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210 <211> LENGTH: 982
211 <212> TYPE: DNA
212 <213> ORGANISM: hepatitis B virus
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                                                                      120
219 ttcaggaaca gtaaaccctg ttccgacttc tgtctctcac acatcgtcaa tcttctcgag
                                                                      180
240
223 cgtgttacag gcggggtttt tcttgttgac aagaatcctc acaataccgc agagtctaga
                                                                      300
225 ctcgtggtgg acttctctca attttctagg gggaactacc gtgtgtcttg gccaaaattc
                                                                      360
227 gcagtcccca acctccaatc actcaccaac ctcctgtcct ccaacttgtc ctggttatcg
                                                                      420
229 ctggatgtat ctgcggcgtt ttatcatctt cctcttcatc ctgctgctat gcctcatctt
                                                                      480
231 cttgttggtt cttctggact atcaaggtat gttgcccgtt tgtcctctaa ttccaggatc
                                                                      540
                                                                      600
233 ttcaaccacc agcacgggac catgcagaac ctgcacgact cctgctcaag gaaactctat
                                                                      660
235 gtatecetee tgttgetgta ceaaacette ggaeggaaat tgeacetgta tteccatece
237 atcatectqq qettteqqaa aatteetatq qqaqtqqqee teaqeeeqtt teteetgget
                                                                      720
239 cagtttacta gtgccatttg ttcagtggtt cgtagggctt tcccccactg tttggctttc
                                                                      780
241 agttatatgg atgatgtggt attgggggcc aagtctgtat cgcatcttga gtcccttttt
                                                                      840
                                                                      900
245 agatggggtc actetttaca tttcatgggc tatgtcattg gatgttatgg gtcattgcca
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247 caagatcaca tcagacagaa aa
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251 <211> LENGTH: 268
252 <212> TYPE: PRT
253 <213> ORGANISM: hepatitis B virus
255 <400> SEQUENCE: 5
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261 Pro Arg Thr Pro Ala Arg Val Thr Gly Gly Val Phe Leu Val Asp Lys
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TIME: 14:53:50

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Output Set: N:\CRF4\02152007\J576906.raw

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262	_			20			~ 1		25	.	**- 7	**- 7	3	30	0	a1
	Asn	Pro		Asn	Thr	Ата	Glu		Arg	Leu	vai	vai		Pne	ser	GIII
266		_	35		_	_	_	40	_	_	_	_	45			_
	Phe		Arg	GLY	Asn	Tyr	Arg	Val	Ser	Trp	Pro		Pne	Ala	Val	Pro
270		50	_				55					60				
273	Asn	Leu	Gln	Ser	Leu	Thr	Asn	Leu	Leu	Ser		Asn	Leu	Ser	\mathtt{Trp}	
274						70					75					80
277	Ser	Leu	Asp	Val	Ser	Ala	Ala	Phe	Tyr	His	Leu	Pro	Leu	His	Pro	Ala
278					85					90					95	
281	Ala	Met	${\tt Pro}$	His	Leu	Leu	Val	Gly	Ser	Ser	Gly	Leu	Ser	Arg	Tyr	Val
282				100					105					110		
285	Ala	Arg	Leu	Ser	Ser	Asn	Ser	Arg	Ile	Phe	Asn	His	Gln	His	Gly	Thr
286		_	115					120					125			
289	Met	Gln	Asn	Leu	His	Asp	Ser	Cys	Ser	Arg	Lys	Leu	Tyr	Val	Ser	Leu
290		130				_	135	•		_	-	140	-			
293	Leu	Leu	Leu	Tyr	Gln	Thr	Phe	Gly	Arq	Lys	Leu	His	Leu	Tyr	ser	His
	145			•		150		-	_	•	155			-		160
		Ile	Ile	Leu	Glv		Arg	Lvs	Ile	Pro		Glv	Val	Glv	Leu	Ser
298					165		5		_	170		-		-	175	
	Pro	Phe	Leu	Len		Gln	Phe	Thr	Ser		Ile	Cvs	Ser	Val	Val	Arg
302				180					185			-1		190		3
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306	9		195			-7-		200			-1-		205			
	Leu	Glv		Lvs	Ser	Val	Ser		Len	Glu	Ser	Leu		Thr	Ala	Val
310		210			-		215					220				
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318	_, _			0 -1	245					250	- 2	-1-			255	-1-
	Tvr	Glv	Ser	Leu		Gln	Asp	His	Ile		Gln	Lvs				
322	-1-	0-1		260			÷E		265			-1 -				
	<210)> SI	O TI		. 6								•			
		l> LI														
		2> T														
	<213> ORGANISM: hepatitis B virus															
)> SI			— ·											
						Ser	Gly	Phe	Leu	Glv	Pro	Leu	Leu	Val	Leu	Gln
333					5		1			10					15	
		Glv	Phe	Phe	_	Leu	Thr	Ara	Ile		Thr	Ile	Pro	Gln		Leu
337		-		20				_	25					30		
	Asp					Ser	Leu				Glv	Glv	Thr			Cvs
341	ASP	DCI	35	111	1111	DCI	שטע	40	1110			017	45			O _I S
	Lau	Glv		λen	Car	Gln	Ser		Thr	Sar	Δen	Hic		Pro	Thr	Ser
345	neu	50	3111	won	JUL	9111	55	110	1111	JGI	11011	60	JUL	110		
	Cvc		Dro	Thr	Cve	Dro	Gly	ጥኒ፣ም	Δνα	מאנו	Met		T.e.11	Δτα	Δτα	Phe
	_	FIO	FIO	TIIL	Cys	70	GTA	TÄT	AT 9	тър	75	TÀT	⊥eu.	nry.	n. y	80
349		т1 ~	Dha	T 011	Dha		Len	T 611	T 011	Ctra		Tla	Dho	Lou	T.e.u	
						TIE	⊔eu	LEU	படப	LVS	படப	115	FILE	TIC II	ueu	٧al
	Ile	116	rne	11Cu						_						
353					85		Met			90					95	

RAW SEQUENCE LISTING DATE: 02/20/2007 PATENT APPLICATION: US/10/576,906 TIME: 14:53:50

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Output Set: N:\CRF4\02152007\J576906.raw

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                                    105
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364 Gln Gly Asn Ser Met Tyr Pro Ser Cys Cys Cys Thr Lys Pro Ser Asp
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368 Gly Asn Cys Thr Cys Ile Pro Ile Pro Ser Ser Trp Ala Phe Gly Lys
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372 Phe Leu Trp Glu Trp Ala Ser Ala Arg Phe Ser Trp Leu Ser Leu Leu
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                    165
376 Val Pro Phe Val Gln Trp Phe Val Gly Leu Ser Pro Thr Val Trp Leu
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                                    185
                                                         190
377
380 Ser Val Ile Trp Met Met Trp Tyr Trp Gly Pro Ser Leu Tyr Arg Ile
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                                                     205
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394 <211> LENGTH: 1045
395 <212> TYPE: DNA
396 <213> ORGANISM: hepatitis B virus
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403 aagetetget agateeeaga gtgaggggee tatattttee tgetggtgge teeagtteeg
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405 gaacagtaaa ccctgttccg actactgcct ctcccatatc gtcaatcttc tcgaggactg
                                                                          240
407 gggaccetge acegaacatg gagageacea cateaggatt cetaggacee etgetegegt
                                                                          300
409 tacaggcggg gtttttcttg ttgacaagaa tcctcacaat accacagagt ctagactcgt
                                                                           360
                                                                           420
411 ggtggacttc tctcaatttt ctagggggaa cacccaagtg tcctggccaa aatttgcagt
413 ccccaacctc caatcactca ccaacctctt gtcctccaat ttgtcctggt tatcgctgga
                                                                           480
415 tgtgtctgcg gcgttttatc atcttcctct tcatcctgct gctatgcctc atcttcttgt
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417 ggggtcttct ggactaccaa ggtatgttgc ccgtttgtcc tctacttcca ggaacatcaa
                                                                           600
                                                                          660
419 ctaccagcac gggaccatgc aagacctgca cgactcctgc tcaaggaacc tctatgtttc
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421 cctcttgttg ctgtacaaaa ccttcggacg gaaattgcac ttgtattccc atcccatcat
423 cttgggcttt cgcaagattc ctatgggagt gggcctcagt ccgtttctcc tggctcagtt
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425 tactagtgcc atttgttcag tggttcgtag ggctttcccc cactgtttgg cttttagtta
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427 tatggatgat gtggtattgg gggccaagtc tgtacaacay cttgaatccc tttttaccgc
                                                                          900
429 tgttaccaat tttcttttgt ctttgggtat acatttaaac cctactaaaa ccaaacgttg
                                                                          960
                                                                          1020
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436 <210> SEQ ID NO: 8
437 <211> LENGTH: 271
438 <212> TYPE: PRT
439 <213> ORGANISM: hepatitis B virus
441 <400> SEQUENCE: 8
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447 Pro Arg Thr Pro Ala Arg Val Thr Gly Gly Val Phe Leu Val Asp Lys
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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/576,906

DATE: 02/20/2007 TIME: 14:53:51

Input Set : A:\19781.seq.txt

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:9; Xaa Pos. 208
Seq#:12; Xaa Pos. 172
Seq#:14; Xaa Pos. 288
Seq#:15; Xaa Pos. 298,300
Seq#:27; Xaa Pos. 160
Seq#:30; Xaa Pos. 95
Seq#:32; Xaa Pos. 312
Seq#:33; Xaa Pos. 309,327
Seq#:38; Xaa Pos. 115,147,239
Seq#:39; Xaa Pos. 134,148,240
Seq#:41; Xaa Pos. 247
Seq#:42; Xaa Pos. 142,247

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/576,906

DATE: 02/20/2007 TIME: 14:53:51

Input Set : A:\19781.seq.txt

Output Set: N:\CRF4\02152007\J576906.raw

L:572 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:192
L:755 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:160
L:895 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:272
L:997 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:288
L:1689 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:144
L:1844 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:80
L:2016 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:304
L:2122 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:304
M:341 Repeated in SeqNo=33
L:2405 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:112
M:341 Repeated in SeqNo=38
L:2508 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:128
M:341 Repeated in SeqNo=39
L:2656 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:240
L:2722 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42 after pos.:128
M:341 Repeated in SeqNo=42